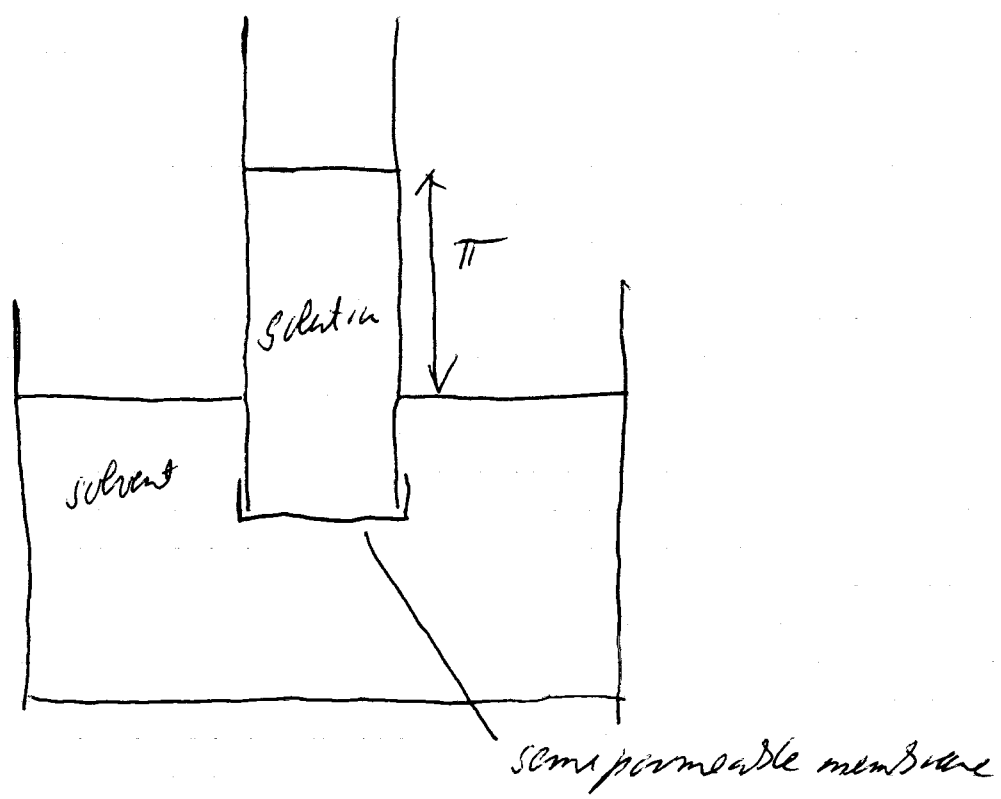


Osmosis



$$\left\{ \begin{aligned} \mu_A^*(p) &= \mu_A(x_A, p+\pi) \\ \mu_A(x_A, p+\pi) &= \mu_A^*(p+\pi) + RT \ln x_A \quad (\text{ideal}) \\ \mu_A^*(p+\pi) &= \mu_A^*(p) + \int_p^{p+\pi} V_m dp \end{aligned} \right.$$

$$\rightarrow -RT \ln x_A = \int_p^{p+\pi} V_m dp$$

define: $\ln x_A = \ln(1-x_B) \approx -x_B$, V_m p-independent

$$RT x_B = \pi V_m \quad x_B \approx \frac{n_B}{n_A} \quad n_A V_m = V$$

$$\Rightarrow \boxed{\pi = \frac{n_B}{V} RT} \quad \text{van't Hoff equation}$$

Binding of small solutes to proteins

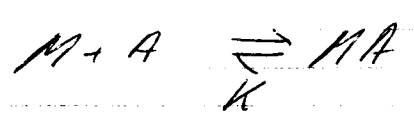
concentration of a protein or other macromolecule (= solute which cannot pass membrane). $[M]$

small solute A which can pass membrane and binds to M: $[A_{in}] = [A_{free}] + [A_{bound}]$

$[A_{free}] = [A_{out}]$ - can be measured

⇒ the average number of A molecules bound to M:

$$\theta = \frac{[A_{bound}]}{[M]} = \frac{[A_{in}] - [A_{out}]}{[M]} \leftarrow \text{measured}$$



$$K = \frac{[MA]}{[M_{free}][A_{free}]} = \frac{[A_{bound}]}{([M] - [A_{bound}])([A_{free}])}$$

~~No dissociation and independent binding sites~~

$$K = \frac{\theta}{(1 - \theta)[A_{out}]}$$

~~Measurement of molecular masses of macromolecules:~~

Measurement of molecular masses of macromolecules:

for non-ideal solution

$$\pi = RT \left(\frac{n_B}{V} + B \frac{n_B^2}{V^2} \right)$$

$$\pi = \rho g h, \quad \frac{n_B}{V} = \frac{c}{M} \quad c \left[\frac{\text{mg}}{\text{ml}} \right]$$

$$\begin{aligned} \Rightarrow \frac{h}{c} &= \frac{RT}{\rho g M} \left(1 + B \frac{c}{M} + \dots \right) = \\ &= \frac{RT}{\rho g M} + \left(\frac{RTB}{\rho g M^2} \right) c + \dots \end{aligned}$$

⇒ Plot h/c against c ⇒
straight line intercepts for $c=0$
at $\frac{RT}{\rho g M} \Rightarrow M$ determined

